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Sequence Listing was accepted.
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Reviewer: Anne Corrigan
Timestamp: [year=2008; month=4; day=24; hr=15; min=8; sec=10; ms=375;]
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Validated By CRFValidator v 1.0.3

Application No: 10091912 Version No: 2.0

Input Set:

Output Set:

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No. of SeqIDs Defined:	3
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<110> Bott, Richard R.
Kellis, James T.
Morrison, Thomas B.

<120> High Throughput Mutagenesis Screening
Method

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<140> 10091912
<141> 2002-03-05

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<170> FastSEQ for Windows Version 4.0

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35 40 45
Arg Ile Tyr Arg Pro Arg Asp Leu Gly Gln Gly Gly Val Arg His Pro
50 55 60
Val Ile Leu Trp Gly Asn Gly Thr Gly Ala Gly Pro Ser Thr Tyr Ala
65 70 75 80

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Gly Leu Leu Ser His Trp Ala Ser His Gly Phe Val Val Ala Ala Ala
                85                  90                  95
Glu Thr Ser Asn Ala Gly Thr Gly Arg Glu Met Leu Ala Cys Leu Asp
            100                 105                 110
Tyr Leu Val Arg Glu Asn Asp Thr Pro Tyr Gly Thr Tyr Ser Gly Lys
        115                 120                 125
Leu Asn Thr Gly Arg Val Gly Thr Ser Gly His Ser Gln Gly Gly Gly
    130                 135                 140
Gly Ser Ile Met Ala Gly Gln Asp Thr Arg Val Arg Thr Thr Ala Pro
145                 150                 155                 160
Ile Gln Pro Tyr Thr Leu Gly Leu Gly His Asp Ser Ala Ser Gln Arg
                165                 170                 175
Arg Gln Gln Gly Pro Met Phe Leu Met Ser Gly Gly Gly Asp Thr Ile
            180                 185                 190
Ala Phe Pro Tyr Leu Asn Ala Gln Pro Val Tyr Arg Arg Ala Asn Val
        195                 200                 205
Pro Val Phe Trp Gly Glu Arg Arg Tyr Val Ser His Phe Glu Pro Val
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Gly Ser Gly Gly Ala Tyr Arg Gly Pro Ser Thr Ala Trp Phe Arg Phe
225                 230                 235                 240
Gln Leu Met Asp Asp Gln Asp Ala Arg Ala Thr Phe Tyr Gly Ala Gln
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    50                  55                  60
Tyr Ala Gly Leu Leu Ser His Trp Ala Ser His Gly Phe Val Val Ala
65                  70                  75                  80
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                85                  90                  95
Leu Asp Tyr Leu Val Arg Glu Asn Asp Thr Pro Tyr Gly Thr Tyr Ser
            100                 105                 110
Gly Lys Leu Asn Thr Gly Arg Val Gly Thr Ser Gly His Ser Gln Gly
        115                 120                 125
Gly Gly Gly Ser Ile Met Ala Gly Gln Asp Thr Arg Val Arg Thr Thr
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Ala Pro Ile Gln Pro Tyr Thr Leu Gly Leu Gly His Asp Ser Ala Ser
145                 150                 155                 160
Gln Arg Arg Gln Gln Gly Pro Met Phe Leu Met Ser Gly Gly Gly Asp
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Thr Ile Ala Phe Pro Tyr Leu Asn Ala Gln Pro Val Tyr Arg Arg Ala
            180                 185                 190
Asn Val Pro Val Phe Trp Gly Glu Arg Arg Tyr Val Ser His Phe Glu
        195                 200                 205
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Pro Val Gly Ser Gly Gly Ala Tyr Arg Gly Pro Ser Thr Ala Trp Phe
    210                 215                 220
Arg Phe Gln Leu Met Asp Asp Gln Asp Ala Arg Ala Thr Phe Tyr Gly
225                 230                 235                 240
Ala Gln Cys Ser Leu Cys Thr Ser Leu Leu Trp Ser Val Glu Arg Arg
                245                 250                 255
Gly Leu
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